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NEWS 24 DEC 17 DGENE now includes more than 10 million sequences
NEWS 25 DEC 17 TOXCENTER enhanced with 2008 MeSH vocabulary in
MEDLINE segment
NEWS 26 DEC 17 MEDLINE and LMEDLINE updated with 2008 MeSH
vocabulary
NEWS 27 DEC 17 CA/CAplus enhanced with new custom IPC display
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NEWS EXPRESS 19 SEPTEMBER 2007: CURRENT WINDOWS VERSION IS V8.2,
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=> s (protease or peptidase or proteinase) (4A) (fusion protein)
L1 776 (PROTEASE OR PEPTIDASE OR PROTEINASE) (4A) (FUSION
PROTEIN)

=> s (protease or peptidase or proteinase) (4A) (auto or self)
L2 571 (PROTEASE OR PEPTIDASE OR PROTEINASE) (4A) (AUTO OR
SELF)

=> s (fusion protein) (4A) (auto or self)
L3 156 (FUSION PROTEIN) (4A) (AUTO OR SELF)

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L4 5 L1 AND L2 AND L3

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=> d 15 1-5 bib ab

L5 ANSWER 1 OF 5 CAPLUS COPYRIGHT 2007 ACS on STN
AN 2005:277491 CAPLUS
DN 142:458829
TI Self-cleavage of fusion protein in vivo
using TEV protease to yield native protein
AU Shih, Yan-Ping; Wu, Hui-Chung; Hu, Su-Ming; Wang, Ting-Fang;
Wang, Andrew
H.-J.
CS Institute of Biological Chemistry and National Core Facilities
of High
Throughput Protein Production, Academia Sinica, Taipei, 115,
Taiwan
SO Protein Science (2005), 14(4), 936-941
CODEN: PRCIEI; ISSN: 0961-8368
PB Cold Spring Harbor Laboratory Press
DT Journal
LA English
AB Overprodn. of proteins from cloned genes using fusion protein
expression
vectors in Escherichia coli and eukaryotic cells has increased
the
quantity of protein produced. This approach has been widely
used in
producing soluble recombinant proteins for structural and
functional anal.
One major disadvantage, however, of applying this approach for
clin. or

bioindustrial uses is that proteolytic removal of the fusion carrier is

tedious, expensive, and often results in products with addnl. amino acid

residues than the native proteins. Here we describe a new method for

productions of native proteins with original amino termini in vivo via

intracellular self-cleavage of the fusion protein using tobacco etch virus (TEV) protease. Our design allows one to simultaneously clone any gene into multiple fusion protein

vectors using two unique cloning sites (i.e., SnaBI and XhoI) without

restriction digestion, and then rapidly identifies those constructs

producing soluble native proteins. This method will make the fusion protein

approach more feasible for protein drug research.

RE.CNT 12 THERE ARE 12 CITED REFERENCES AVAILABLE FOR THIS RECORD
ALL CITATIONS AVAILABLE IN THE RE FORMAT

L5 ANSWER 2 OF 5 CAPLUS COPYRIGHT 2007 ACS on STN

AN 2003:875431 CAPLUS

DN 139:359898

TI Reporter-selectable hepatitis C virus replicon and its stably transfected

hepatoma cell line for drug screening

IN Duggal, Rohit; Patick, Amy Karen; Zhang, Jie; Zhao, Weidong

PA Pfizer Inc., USA

SO PCT Int. Appl., 52 pp.

CODEN: PIXXD2

DT Patent

LA English

FAN.CNT 1

| PATENT NO. | KIND | DATE | APPLICATION NO. |
|------------|------|------|-----------------|
| DATE | | | |

PI WO 2003091439 A1 20031106 WO 2003-IB1687
20030422

W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA,
CH, CN,
CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD,
GE, GH,
GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC,
LK, LR,
LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NI, NO,
NZ, OM,
PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, TJ, TM, TN,
TR, TT,

TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW
RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AM,
AZ, BY, KG, KZ, MD, RU, TJ, TM, AT, BE, BG, CH, CY, CZ, DE, DK,
EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI,
SK, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN,
TD, TG
CA 2483513 A1 20031106 CA 2003-2483513
20030422 AU 2003225479 A1 20031110 AU 2003-225479
20030422 EP 1499727 A1 20050126 EP 2003-747188
20030422 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE,
MC, PT, IE, SI, LT, LV, FI, RO, MK, CY, AL, TR, BG, CZ, EE, HU,
SK JP 2005523705 T 20050811 JP 2003-587967
20030422 MX 2004PA10548 A 20050125 MX 2004-PA10548
20041025 PRAI US 2002-375667P P 20020426
WO 2003-IB1687 W 20030422

AB The invention relates to a reporter-selectable hepatitis C virus (HCV) replicon, and use of the replicon to generate stable, human hepatoma cell

lines. Specifically, a replicon (BB7-M4-hRLuc) has been constructed

containing the 5' NTR fused to a small portion of the core coding region, the

humanized Renilla luciferase gene (hRLuc), a self-cleaving peptide of foot

and mouth disease virus (FMDV) 2A proteinase, the NPTII gene, and an EMCV

IRES (designated EI), followed by the NS3 to NS5B HCV coding region and

the 3' NTR region. The replicon has two adaptive mutations in NS3 (E1202G

and T1280I) and one in NS5A (S2197P). The stable hepatoma cell line

(BB7M4hRLuc#10) stably transfected with this HCV replicon is capable of

generating 700,000 relative light units (RLU, units for expressing

luciferase activity) of reporter gene activity, which amts. up to a 70

fold difference in signal to noise ratios compared to other available

stable cell line. The replicon and cell lines are useful in the compound screening process in HCV drug discovery.

RE.CNT 5 THERE ARE 5 CITED REFERENCES AVAILABLE FOR THIS RECORD
ALL CITATIONS AVAILABLE IN THE RE FORMAT

L5 ANSWER 3 OF 5 CAPLUS COPYRIGHT 2007 ACS on STN
AN 1996:510609 CAPLUS
DN 125:161813
TI Cloning, expression and purification of HIV-1 protease
AU Wan, Min; Takagi, Masahiro
CS Faculty Medicine, National University Singapore, Singapore
SO Microbial Utilization of Renewable Resources (1996), Volume Date
1995, 9,
348-361

CODEN: MURRE6

PB International Center for Cooperative Research in Biotechnology,
Japan

DT Journal

LA English

AB Various constructs containing the N-terminal extended human immunodeficiency

virus type 1 (HIV-1) protease gene (PR 107) were studied. The constructs

of PR 107 gene expressed as fusion protein with the glutathione S-transferase of the GST gene or the maltose-binding protein (MBP) of the

Mal E gene showed that the full-length fusion protein exhibited self-processing in E. coli. The results from expts. indicated that the size of the fusion portion could not affect the

self-processing of HIV-1 protease obviously. Although the larger fusion portion (MBP) may offer bigger steric-interference for

the formation of the appropriate conformation of the fused protease and therefore lower the self-cleavage process, the protease could still easily self-process from the fusion portion to release itself, despite that only one subunit of the

dimeric protease attached to GST or MBP. An isolation method consisting

of denaturation of protein and followed by refolding was developed for

releasing this recombinant HIV-1 PR into the soluble phase since most of the

expressed protease was initially present in insol. inclusion bodies. More

than 600-fold purification was obtained by sequential purification using Sephadex

G-50 gel filtration and CM-23 cellulose cation exchange chromatog.,

yielding the protease whose purity was more than 95%. SDS-PAGE indicated

that the mol. weight of this recombinant HIV-1 PR is 11 kDa. The recombinant

HIV-1 protease showed proteolytic activity for the synthetic peptide

substrate corresponding to the sequence of gag MA/CA and pol p6*/PR

junctions. Immuno-blotting indicated that these recombinant HIV-1

protease specifically reacted with HIV-1 protease antisera. The purified

enzyme whose specific activity for the heptapeptide SQNYPIV was 848.7

nmol*min-1*mg protease-1 also processed recombinant polyprotein Gag41 as

its substrate.

L5 ANSWER 4 OF 5 CAPLUS COPYRIGHT 2007 ACS on STN

AN 1995:642290 CAPLUS

DN 123:219557

TI Replicating foamy virus-based vectors directing high level expression of

foreign genes

AU Schmidt, Michael; Rethwilm, Axel

CS Institut fuer Virologie und Immunbiologie, Universitaet Wuerzburg,

Wuerzburg, 97078, Germany

SO Virology (1995), 210(1), 167-78

CODEN: VIRLAX; ISSN: 0042-6822

PB Academic

DT Journal

LA English

AB Replication-competent retroviral vectors (pFOV-1 to -3 and -7) were

constructed on the basis of an infectious human foamy virus mol. clone

which has deletions in the U3 region of the long terminal repeat and in

the 3' region of the genome, previously identified to be nonessential for

virus replication in vitro. The CAT and luciferase indicator genes were

expressed as C-terminal fusion proteins to 215 amino acids of the viral

Bet protein in the pFOV-1 vector. Introduction of the foot-and-mouth

disease 2A protease sequence between the truncated bet coding sequence and

the cloning site for the insertion of foreign genes in the pFOV-7 vector

resulted in self-cleaving of the recombinant fusion protein. Alternatively, an internal ribosomal binding site was introduced, allowing expression of authentic foreign protein (pFOV-2 and -3 vectors). DNA fragments derived from the mouse hepatitis virus surface

gene up to the length of 1.3 kb were inserted into pFOV-1. The vector

constructs gave rise to viruses which were fully infectious in diploid

human fibroblasts and recombinant viruses stably expressed high levels of

foreign protein indicating that the pFOV vectors may be useful tools to

study the effects of proteins of interest at least in tissue culture

cells.

L5 ANSWER 5 OF 5 CAPLUS COPYRIGHT 2007 ACS on STN

AN 1991:576747 CAPLUS

DN 115:176747

TI Self-cleaving fusion proteins

IN Louis, John M.

PA National Institutes of Health, USA

SO U. S. Pat. Appl., 38 pp. Avail. NTIS Order No. PAT-APPL-6-586
079.

CODEN: XAXXAV

DT Patent

LA English

FAN.CNT 1

| | PATENT NO. | KIND | DATE | APPLICATION NO. |
|--------------------------|------------|----------|----------------|-----------------|
| DATE | | | | |
| ----- | ----- | ----- | ----- | ----- |
| PI US 586079
19900921 | A0 | 19910801 | US 1990-586079 | |
| US 6077694 | A | 20000620 | | |
| WO 9205276
19910920 | A1 | 19920402 | WO 1991-US6735 | |

W: AU, CA, JP

RW: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LU, NL, SE

AU 9185377 A 19920415 AU 1991-85377

19910920

PRAI US 1990-586079 A 19900921
WO 1991-US6735 A 19910920

AB Self-processing fusion proteins are used for the manufacture of a protein of

interest. The fusion protein has three domains: an affinity domain for

rapid purification of the protein by affinity chromatog.; a retroviral

proteinase domain flanked by cleavage sites recognized by the proteinase;

and, a domain of interest (the target protein). After purification of the

protein by affinity chromatog. the protein is partially denatured allowing

the proteinase to cleave it into three domains. When the fusion protein

is accumulated as inclusion bodies, it may be cleaved during the denaturation/solubilization process. Use of sequences from the pol region

of the retrovirus gene can be used to control solubility of the protein. The

use of this procedure to purify and accurately cleave soluble and insol.

fusion proteins of human immunodeficiency virus proteinase and the

Escherichia coli malE gene product was demonstrated.

EAST Search History

| Ref # | Hits | Search Query | DBs | Default Operator | Plurals | Time Stamp |
|-------|------|---------------------------------|--------------------|------------------|---------|------------------|
| L1 | 1039 | protease near4 (fusion protein) | US-PGPUB;
USPAT | ADJ | OFF | 2007/12/20 20:10 |
| L2 | 83 | protease near4 (auto) | US-PGPUB;
USPAT | ADJ | OFF | 2007/12/20 20:11 |
| L3 | 6 | I1 and I2 | US-PGPUB;
USPAT | ADJ | OFF | 2007/12/20 20:11 |

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